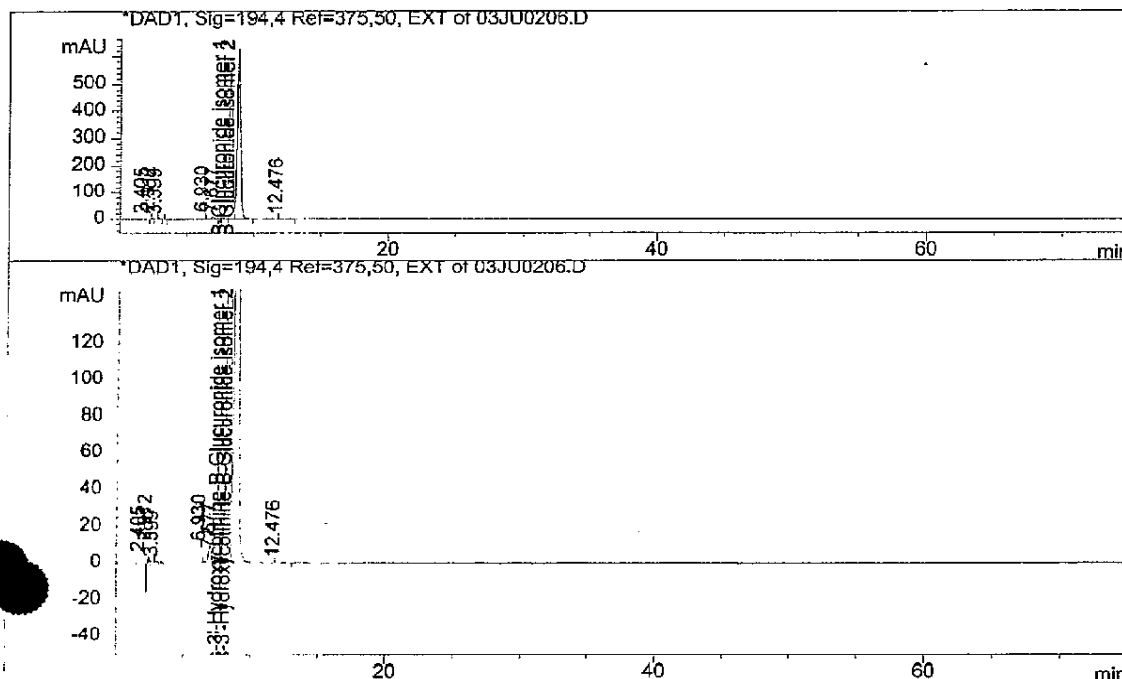


Injection Date : Mon, 3. Jun. 2002 Seq Line : 2
 Injection Time : 11:08:08 PM Vial No. : 91
 Sample Name : 3338.A* Inj. No. : 4
 Acq Operator : admin
 Acq. Method : PURITY2.M
 Analysis Method : C:\MDSPS\PROJECT\3338A\3338A_P1.M
 Last Changed : Tue, 4. Jun. 2002, 09:59:13 am

HPLC PURITY PROFILING OF MDSPSR_3338.A*



Customized Report: HPLC PROFILING

Sorted By Signal
 Calib. Data Modified : Tue, 4. Jun. 2002, 09:58:52 am
 Multiplier : 1.000000
 Dilution : 1.000000
 Uncalibrated Peaks : not reported

Signal Description : DAD1, Sig=194,4 Ref=375,50, EXT

#	RT	Type	Height	Area	Area %	Name
---	[min]		-----	-----	-----	
1	2.405	BBA	2.809	12.6097	0.064	
2	2.912	BBA	8.947	46.5298	0.237	
3	3.399	PBA	0.758	2.7246	0.014	
4	6.930	BP	9.149	133.5809	0.681	
5	7.677	VV	4.954	58.9878	0.301	
6	8.257	VV	562.666	8884.6650	45.307	trans-3'-Hydroxyc
7	8.824	VB	629.477	10463.3574	53.358	trans-3'-Hydroxyc
8	12.476	VP	0.238	7.3603	0.038	

Totals:

*** End of Report ***

Average Purity Profiling of:
 MDSPSR_3338.A* isomer 1 = 45.3
 MDSPSR_3338.A* isomer 2 = 53.4
 MDSPSR_3338.A* (total) = 98.7
 JT 04/JUN/02
 Average Retention time of:
 MDSPSR_3338.A* isomer 1 = 8.27 min
 MDSPSR_3338.A* isomer 2 = 8.84 min
 JT 04/JUN/02